c/C	JAN 2 B ZOTES STATES			
\	JAN 2 B JOHN STEE			
1	ATENT & TRADE			
	475.18 P>3.00	WI-22211		Homo sapiens full length insert cDNA clone
	475.18 P2.08	stSG48093		ESTs
	475.18 *	A004P27		ESTs, Weakly similar to MULTIDRUG RESI
	475.35 P2.10	stSG9904		ESTs
	475.40 P0.45	sts-AA024696		ESTs
	475.51 P>3.00	stSG53793		ESTs
	476.10 P>3.00	Bda98d05		Homo sapiens full length insert cDNA clone
	476.21 P>3.00	sts-H24468		ESTs
1	476.21 P>3.00	sts-N94741		ESTs
1	476.64 P0.28	stSG22488		ESTs
١	476.85 P0.36	stSG44909		ESTs
1	477.06 PO.10	stSG54797		ESTs
1	477.27 P1.33	stSG48099		ESTs
	477.37 P0.09 *	sts-AA028894		Homo sapiens silencing mediator of retinoic
	477.80 P1.44	stSG52727		EST
1	477.80 "	U44799		Human U1-snRNP binding protein homolog mR
ı	477.80 "	WI-15963		ESTs
1	477.80 "	stSG53886		ESTs, Weakly similar to neuronal thread pr
1	478.74 PO,01	WIAF-364		ESTs
1	479.01 P0.21	WI-21080		ESTs
1	479.13 PO.19	A009B29		ESTs
1	479.33 PO.22	A006F32	EIF2B1	eukaryotic translation initiation factor 2B
	479.33 PO.19	WIAF-449	EIF2B1	eukaryotic translation initiation factor 2B
1	479.33 P0.19 *	WI-15890		H.sapiens mRNA for transmembrane protein r
1	479.55 P0.20 *	stSG349		H.sapiens mRNA for transmembrane protein r
	479.55 " *	A004O46	BDKRB2	bradykinin receptor B2
	479.55 "	stSG42540		ESTs
	479.55 "	sts-N26791		ESTs
1	479.55 "	stSG53943		ESTs
	479.55 "	stSG49468		EST
	479.74 P0.16	AFM294ze9	D12S342	Microsatellite marker AFM294ze9
	481.46 P0.00	sts-AA007694		EST
+	=◆ 481.56 F	AFM294xg1	D12S340	Microsatellite anchor marker AFM294xg1
1	. Next interval down			

FIG. 2I

145.7

147.5

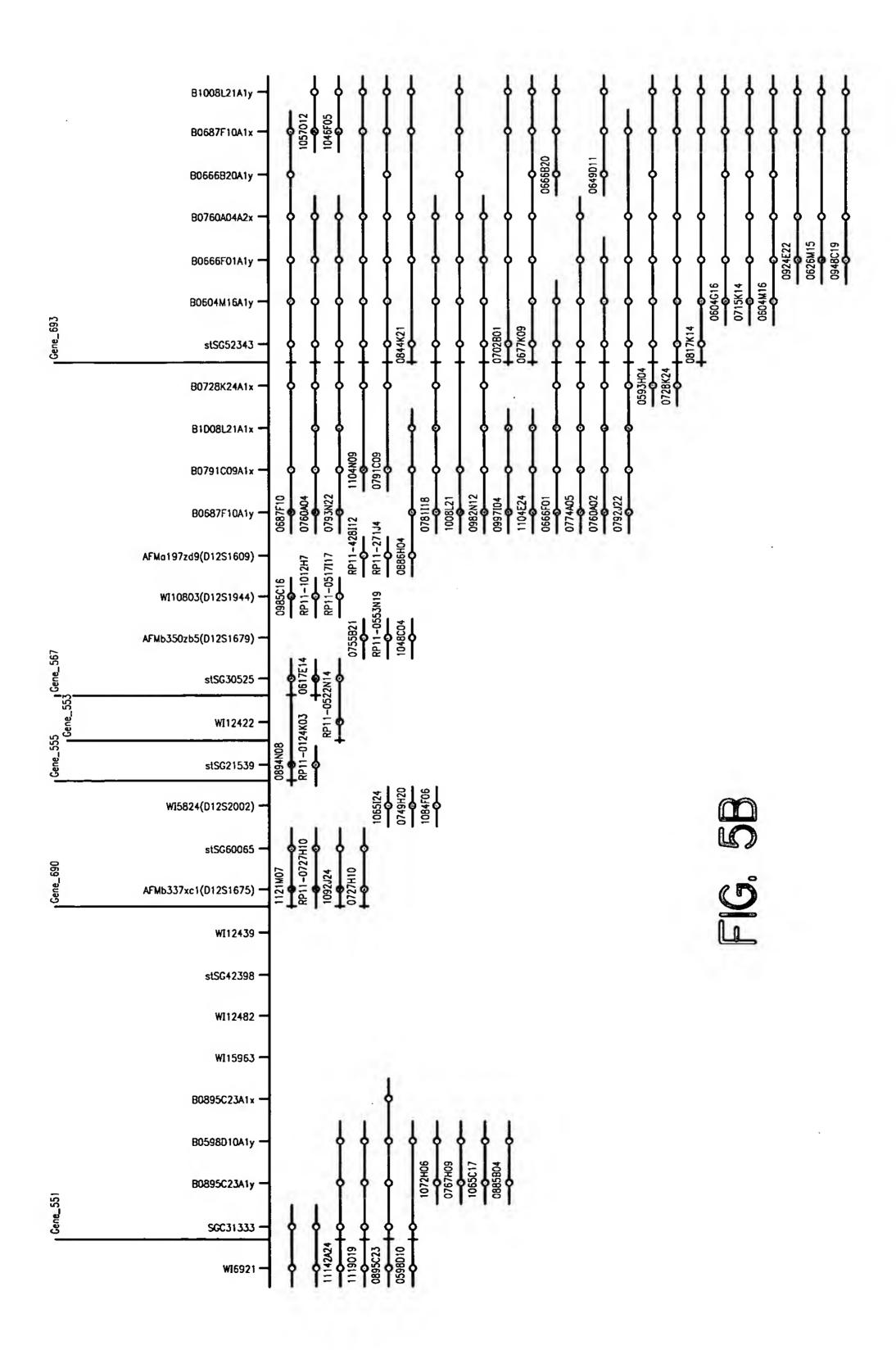
483.87 P0.83		stSG47707		ESTs
484.70 P0.93		stSG4060		ESTs
484.70 "		stSG62390	GTF2H3	general transcription factor IIH, polypepti
484.70 "		stSG42994		ESTs
484.73 .P0.74		stSG46906		ESTs
484.80 P0.91		A004X33		ESTs
484.91 P1.11		stSG3211		ESTs, Weakly similar to B-cell growth fact.
484.91 "	*	sts-Z41302	BDKRB2	bradykinin receptor B2
484.91 "	*	sts-Z41302	BDKRB2	bradykinin receptor B2
484.91 "		sts-T58259		ESTs, Weakly similar to B-cell growth fact
484.91 "		stSG52737		ESTs
484.91 "		Bda03b10	UBC	ubiquitin C
484.91 "		stSG1936	CD36L1	CD36 antigen (collagen type I receptor, thr
484.91 "		sts-AA017225		ESTs
484.91 P1.15		WI-12212		ESTs
485.12 P1.18		A004F14		ESTs
485.12 P1.18		SGC31333		ESTs
485.23 P1.21	*	WI-12482	BDKRB2	bradykinin receptor B2
485.23 P1.07		sts-AA017698		ESTs
485.33 P1.22		WI-12422		ESTs
485.51 P1.18		stSG42398		EST
485.64 P1.04		sts-AA009669		ESTs
486.07 P2.50		stSG21539		EST
486.13 P1.44		WI-12439		EST
486.34 P1.26		sts-W31616	UBC	ubiquitin C
486.38 P>3.00		stSG54715		ESTs
486.76 P1.64	*	WI-6921		H.sapiens mRNA for transmembrane protein r
487.08 P>3.00		WI-13120		Human mRNA for KIAA0318 gene, partial cds
487.23 P>3.00		stSG54353		ESTs
487.23 P>3.00		stSG22703		EST .
487.28 P>3.00		stSG62698		ESTs
487.28 P>3.00	*	sts-D60472		Homo sapiens silencing mediator of retinoic
487.28 P>3.00		stSG36097		ESTs
487.33 P1.36		sts-U37146		Homo sapiens silencing mediator of retinoic

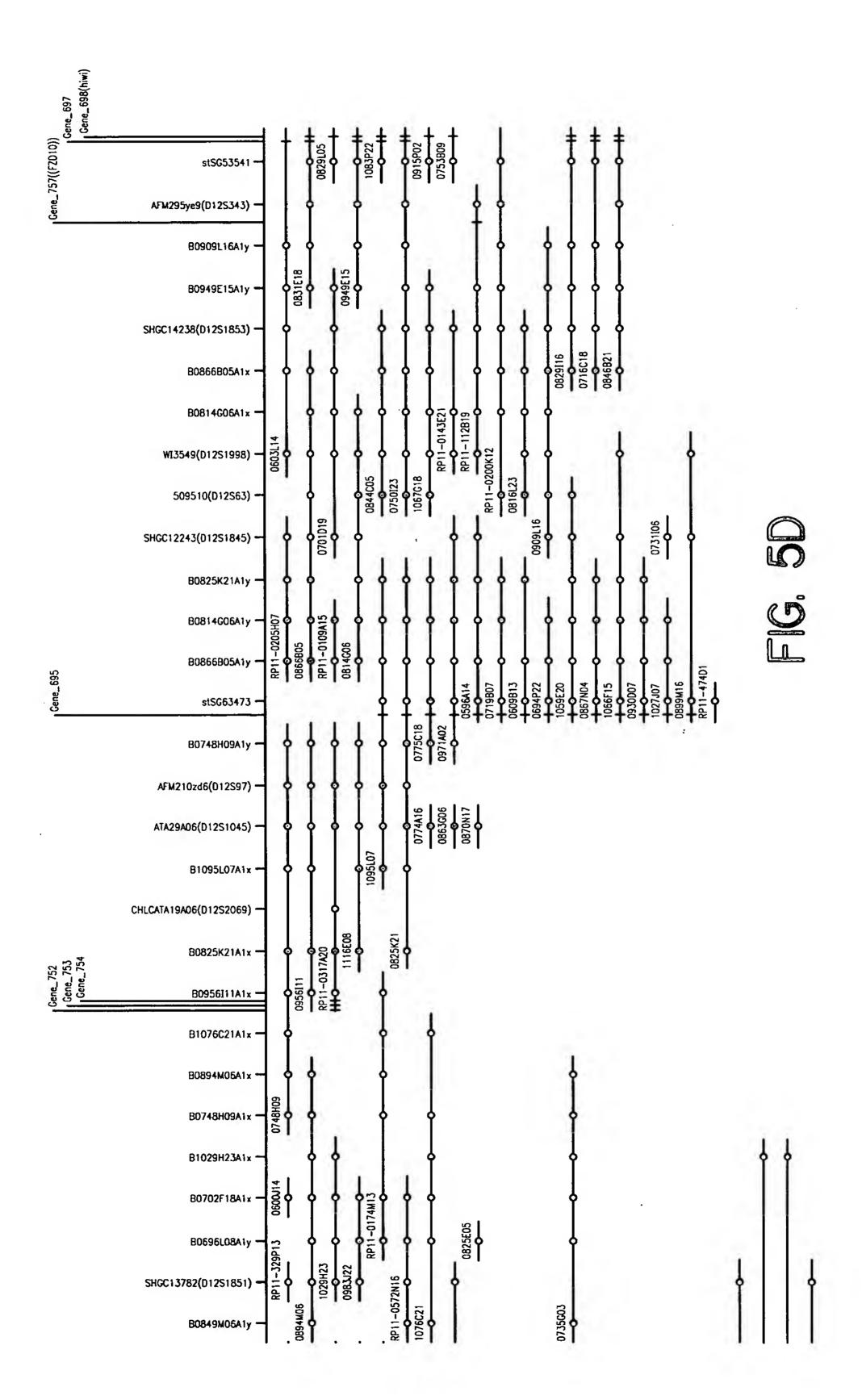
FIG. 2K

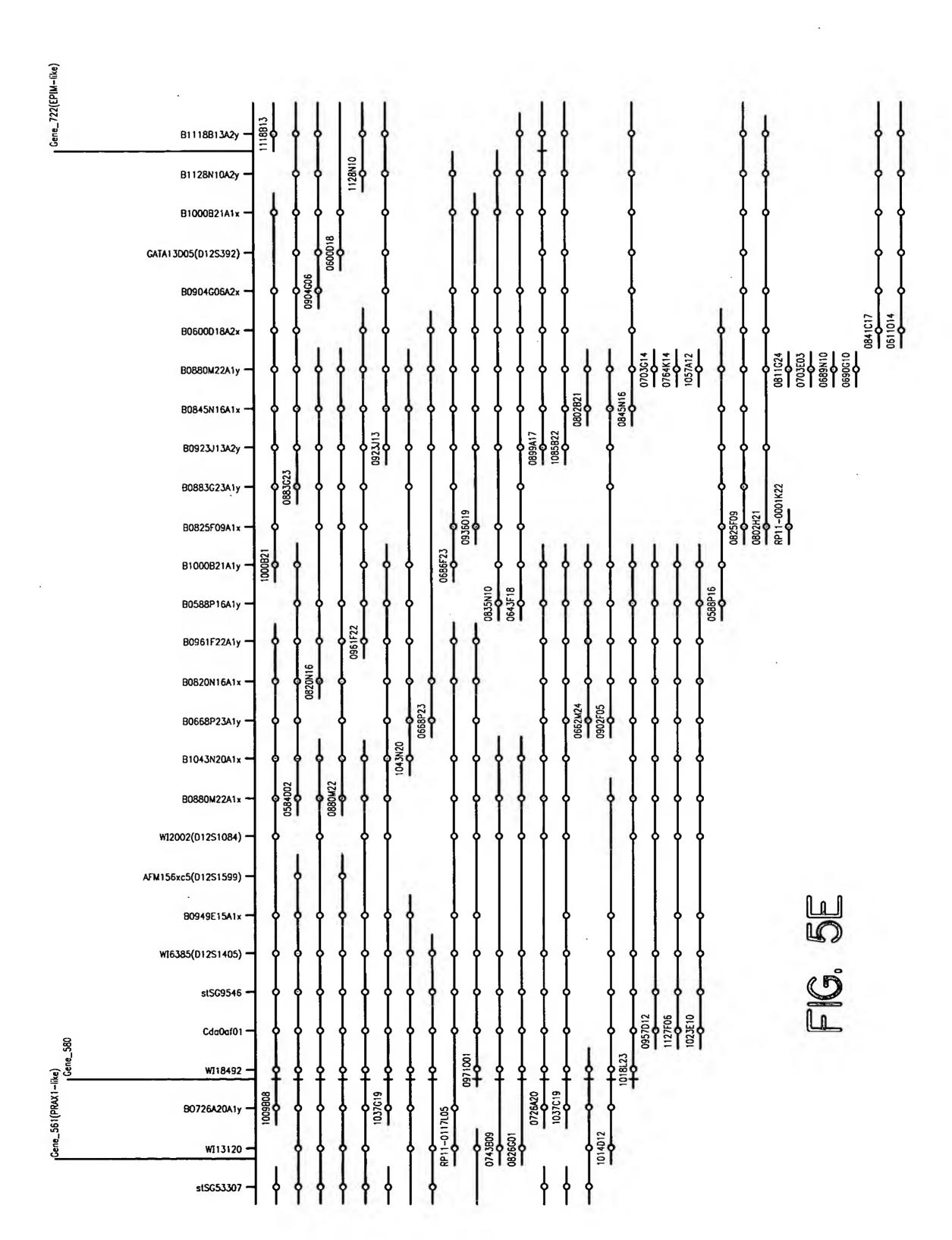
			405.40					
				P>3.00		stSG9807		ESTs
	1		487.50	P>3.00		stSG15434		ESTs
			487.60	P>3.00		stSG53251		ESTs
			487.60	P>3.00		stSG30525	SRRP129	SC35-interacting protein 1
			487.60	P>3.00		stSG46424	D12S1609	ESTs
			487.70	P>3.00		A007A34		ESTs
154.4	.4		487.75	P2.00		AFMa197zd9		Microsatellite marker AFMa197zd9
	1		487.75	P2.02		A006D44		ESTs
			487.80	P>3.00		SGC30248		ESTs, Weakly similar to peptide/histidine
			488.07	P1.68		stSG6320		Homo sapiens clone 24617 mRNA sequence
			488.07	P1.66		stSG6305		Homo sapiens clone 24790 mRNA sequence
			488.07	P0.02		sts-N20163		Homo sapiens full length insert cDNA clone
			488.12	P>3.00		stSG60065		ESTs
			488.12	P>3.00		stSG47723	CPN2 D12S367	ESTs
			488.44	P1.59		stSG3292		Homo sapiens clone 24790 mRNA sequence
			488.44	P0.03		WIAF-856		EST, Weakly similar to reverse transcripta
			488.6 5	P1.54		WI-12272		Homo sapiens clone 24790 mRNA sequence
			488.65	P1.82		stSG52343		ESTs
			488.82	P1.80		stSG16387		carboxypeptidase N, polypeptide 2, 83kD
			488.97	P1.80		SGC31722		ESTs
			489.07	P0.06		stSG54325		ESTs
			489.07	P>3.00		stSG63473		ESTs
160.9	• •	>	489.07	P>3.00		AFMa123xel		Microsatellite marker AFMa123xe1
	1		489.14	P0.17		sts-T81113		ESTs
	-		489.29	P0.05		sts-AA025438		EST
			489.50	P1.37	***	Cdalad08		ESTs
			489.50	P0.05		WI-15018		ESTs
			489.50	P1.50		WI-18492		ESTs
			489.57	P1.48		WI-16177		Homo sapiens androgen receptor associated p
			489.67	P1.44		stSG53307		ESTs
	1		489.71	P1.43		stSG53541		Homo sapiens hiwi mRNA, partial cds
			489.71	P1.43		stSG9546		Homo sapiens clone 24617 mRNA sequence
			489.89	P1.56		A006O16		ESTs
			490.10	P1.42		H64839		EST

FIG. 2L

499.71 P1.72	stSG30906		ESTs
499.71 "	stSG43796	MMP17	matrix metalloproteinase 17 (membrane-insert
499.71 P1.71	sts-X89576	MMP17	matrix metalloproteinase 17 (membrane-insert
499.92 P>3.00	stSG43769		ESTs
500.50 P1.88	stSG26056		ESTs
500.50 P2.33	SGC30786	KIAA0331	KIAA0331 gene product
500.61 P>3.00	stSG1702		Homo sapiens CAGH32 mRNA, partial cds
500.61 *	sts-N59820		ESTs
500.61 "	stSG42115	KIAA0331	KIAA0331 gene product
500.61 "	IB2452	ULK1	unc-51 (C. elegans)-like kinase 1
500.61 "	stSG52521		ESTs
500.61 "	FB9F8		ESTs, Weakly similar to PUTATIVE ATP-D
500.61 "	AA252357		ESTs
500.61 "	stSG4720		Homo sapiens pseudouridine synthase 1 (PUS
500.61 "	sts-AA001424	KIAA0331	KIAA0331 gene product
500.61 P>3.00	stSG31443		ESTs
500.61 P>3.00	stSG49622	ULK1	unc-51 (C. elegans)-like kinase 1
500.61 P2.49	stSG50559		ESTs
501.04 P1.10	stSG54842		ESTs
501.04 P2.03	A008Y05		Unknown
501.89 P2.18	stSG39493		Homo sapiens CAGH32 mRNA, partial cds
501.99 P>3.00	A002A44		Homo sapiens CAGH32 mRNA, partial cds
501.99 P>3.00	sts-H94865	•	EST
501.99 P>3.00	R50113		ESTs
502.10 P1.75	stSG48386		ESTs
502.10 "	stSG50504		ESTs
502.63 P0.06	A006R19		ESTs
502.63 P1.06	WIAF-864		ESTs
502.94 P1.51	stSG54813		ESTs, Weakly similar to peroxisome membran
503.04 P1.42	A004B47		ESTs, Highly similar to DNA polymerase ep
503.25 P0.28	stSG27206		ESTs
503.25 "	stSG40199		Homo sapiens mRNA for KIAA0692 protein, p
503.46 P0.23	stSG8935		ESTs
504.68 P0.69	stSG4731		Homo sapiens mRNA for KIAA0692 protein, p







B0723P10A1y 0714009 0912H06 0619F02 0918D14 0918D14 0582G14 B1056C02A2x stSG54325 stSG42115 B0839D11A1y -987 1 23 4 4 E 4 B1128L12A1y -Gene_751 W16077(D12S1322) -B0716I10A2x SGC34088 -Gene_575 WIAF856 -RP11-0579G13 B0610N03A1x 11281.12 B0700A09A2y RP11-0243M05 B1118B13A2x 1093508 0839011 0909£24 1093H10 B0909E24A1y B0600D18A2y = B1128N10A2x Ŷ B0845N16A1y -B0904G05A2y B0841C17A2x -B0611014A2x -Gene_581(TC4, RAN) stSG3292 -80923J13A2x stSG3357 i Gene_584 i Gene_585 WI16177 H64839 B0700A09A2x W112272 0610N03 0904N07

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